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Role of the receptor for advanced glycation endproducts (RAGE) in retinal vasodegenerative pathology during diabetes in mice

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Abstract

Aims/hypothesis The receptor for AGEs (RAGE) is linked to proinflammatory pathology in a range of tissues. The objective of this study was to assess the potential modulatory role of RAGE in diabetic retinopathy.

Methods Diabetes was induced in wild-type (WT) and *Rage*^{-/-} mice (also known as *Ager*^{-/-} mice) using streptozotocin while non-diabetic control mice received saline. For all groups, blood glucose, HbA_{1c} and retinal levels of methylglyoxal (MG) were evaluated up to 24 weeks post diabetes induction. After mice were killed, retinal glia and microglial activation, vasopermeability, leucostasis and degenerative microvasculature changes were determined.

Results Retinal expression of RAGE in WT diabetic mice was increased after 12 weeks ($p < 0.01$) but not after 24 weeks. *Rage*^{-/-} mice showed comparable diabetes but accumulated less MG and this corresponded to enhanced activity of the MG-detoxifying enzyme glyoxalase I in their retina when compared with WT mice. Diabetic *Rage*^{-/-} mice showed significantly less vasopermeability, leucostasis and microglial activation ($p < 0.05$ – 0.001). *Rage*^{-/-} mice were also protected

against diabetes-related retinal acellular capillary formation ($p < 0.001$) but not against pericyte loss.

Conclusions/interpretation *Rage*^{-/-} in diabetic mice is protective against many retinopathic lesions, especially those related to innate immune responses. Inhibition of RAGE could be a therapeutic option to prevent diabetic retinopathy.

Keywords Acellular capillaries · *Ager* · Diabetic retinopathy · Glucose · Glyoxalase I · HbA_{1c} · Leakage · Methylglyoxal · Microglia · Mouse · *Rage*

Abbreviations

BRB	Blood–retina barrier
Col IV	Collagen IV
GFAP	Glial fibrillary acidic protein
GLO-1	Glyoxalase I
GSH	Glutathione
HMGB-1	High-mobility group box-1
MG	Methylglyoxal
PRR	Pattern-recognition receptor
qPCR	Quantitative RT-PCR
RAGE-Fc	RAGE-inhibiting fusion protein
RAGE	Receptor for AGEs
sRAGE	Soluble protein fragment of RAGE
STZ	Streptozotocin
TUNEL	TdT-mediated dUTP-X nick end labelling
TLR	Toll-like receptor
WT	Wild-type

Dr Angelika Bierhaus died on 15 April 2012, before publication of this work.

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Introduction

Diabetic retinopathy is typified by breakdown of the blood–retina barrier (BRB), loss of capillary pericytes and endothelium, microaneurysm formation, neuronal/glial dysfunction

and progressive ischaemia in the early (non-proliferative) stages [1]. The pathogenesis of diabetic retinopathy is complex and while the underpinning mechanism(s) remains ill-defined, there is strong evidence that inflammation plays an important role [2]. Even in the early stages of disease inflammatory pathology manifests as activation of resident and infiltrating immune cells, abnormal expression of pro-inflammatory cytokines, upregulation of adhesion molecules and leucostasis in retinal capillaries [2]. This relatively non-specific and sustained response to diabetes-related cell injury seems to be driven, at least in part, by microglia and Müller glia acting as the main resident innate immune cells of the retina.

Innate immune responses are regulated by pattern-recognition receptors (PRRs) expressed by immunologically active cells. PRRs, such as the Toll-like receptors (TLRs), CD36 and the receptor for AGEs (RAGE), can orchestrate innate responses such as cytokine release and immune cell activation [3]. Although the complex interplay of PRRs in the diabetic retina remains poorly understood there has been recent attention given to RAGE and its role in diabetic retinopathy [4–6].

Amongst the major ligands for RAGE are high-mobility group box-1 (HMGB-1), amyloid- β , S100B and AGEs [7], many of which occur in the diabetic retina [8, 9]. Inflammation-related pathways involving RAGE are active in retinal endothelial cells [10] and Müller glia [6] and expression of the receptor is enhanced under diabetic conditions [6]. RAGE signalling evokes extracellular signal-regulated kinase 1/2, mitogen-activated protein kinases and P38, leading to downstream activation of nuclear factor- κ B and induction of pro-inflammatory cytokines and/or oxidative stress [11]. Furthermore, blockade of RAGE ligand–receptor binding using a soluble protein fragment (sRAGE) can prevent Müller glial dysfunction [4] during diabetes and retinal capillary leucostasis in AGE-infused normal mice [12]. A recent study by Li et al [5] used a RAGE-inhibiting fusion protein (RAGE-Fc) in diabetic mice and demonstrated a reduced leakage of albumin into neural retina, although there was no significant change in capillary leucostasis. In mice diabetic for 10 months, RAGE blockade prevented acellular capillary formation but not retinal pericyte loss [5]. The current study seeks to investigate retinal lesion progression using mice in which RAGE protein has been deleted and we provide insight into the role RAGE plays in diabetic retinopathy.

Methods

Diabetes induction in mice The *Rage*-knockout mouse (*Rage*^{−/−}; also known as *Ager*^{−/−}) was generated on an SVEV129×C57BL/6 background and backcrossed to C57BL/6 mice for five generations [13]. Wild-type (WT)

C57BL/6 mice were purchased from Harlan Laboratories (Bicester, UK) and maintained within the Biological Research Unit at Queen's University Belfast.

All experiments conformed to the Principles of Laboratory Animal Care (National Institutes of Health, Bethesda, MD, USA) and to the UK Home Office regulations. Diabetes was induced in male *Rage*^{−/−} mice and WT mice at 12 weeks of age using five daily i.p. injections of streptozotocin (STZ) (Sigma, Gillingham, UK) (50 mg/kg in 0.1 mol/l citrate buffer pH 4.6). Control groups received citrate buffer. Hyperglycaemia was confirmed 10 days after injection using glucometric analysis of tail-prick blood samples (FreeStyle Lite; Abbott Laboratories, Dublin, Ireland). Non-fasting blood glucose concentrations >13 mmol/l were considered to indicate diabetes. Blood glucose and weight was monitored bi-weekly. HbA_{1c} levels were determined using Glyco-Tek Affinity Column (Helena Biosciences Europe, Gateshead, UK) whenever mice were killed.

Glyoxalase I activity and methylglyoxal formation in the diabetic retina Glyoxalase I (GLO-1) enzymatic activity in retinal homogenates from the 4-week groups was assayed using previously published protocols [14]. At 8 and 24 weeks diabetes, the synthesis and purification of methylglyoxal (MG), as well as the synthesis of the derivatising agent and the standards for determination of MG, were prepared according to published procedures [14]. The concentration of MG was determined in tissue as nmol/mg and by derivatisation with 1,2-diamino-4,5-dimethoxybenzene. The protein concentration of the tissue homogenate was determined by the Bradford assay using BSA as a standard.

GLO-1 and glutathione immunostaining GLO-1 and glutathione (GSH) immunostaining was assessed using rat anti-GLO-1 (1:200; Abcam, Cambridge, UK) and rabbit anti-GSH (1:1,000; Agrisera, Vännäs, Sweden) on frozen retinal sections (20 μ m). The GLO-1 and GSH were visualised using goat anti-rat antibody labelled with Alexa Fluor 488 and goat anti-rabbit Alexa Fluor 568 (Invitrogen, Paisley, UK).

Quantitative RT-PCR for WT retina Total RNA of retina was extracted using RNeasy Mini Kit (Qiagen, UK). The level of *Rage* and *Glo-1* (*Glo1*) expression in the retina was examined using quantitative RT-PCR (qPCR) as previously reported [15]. β -Actin was used as the housekeeping gene. Primer sequences are listed in electronic supplementary material (ESM) Table 1.

Assessment of the BRB Assessment of the BRB was conducted in cryosections, by assessing albumin leakage from the retinal vasculature. The vasculature was visualised using biotinylated isolectin GS-IB4 (Sigma) followed by binding to streptavidin–Alexa Fluor 488 (Life Technologies, Paisley, UK). Albumin was localised using a goat anti-mouse albumin

antibody (Bethyl Laboratories, Montgomery, TX, USA) followed with anti-goat Alexa Fluor 568 (1:500; Life Technologies). Vessel or neuropile-localised fluorescence was visualised by using a Nikon TE EZ-C1 confocal system (Nikon, Kingston Upon Thames, UK). Using NIS Elements software (Nikon) a threshold value of 12,000 was set for measuring image intensity/brightness. The area (μm^2) of the albumin staining was measured with the 12,000 brightness intensity threshold in the retinal sections. Images were taken at three separate points on the central retina at magnification $\times 40$ and $n=5$ per group of mice.

Leucostasis in the retinal vasculature Leucostasis in diabetic mice was assessed after 4 weeks of diabetes using fluorophore-conjugated concanavalin-A, as described previously [16]. The retinas were mounted on slides and visualised using a Nikon TE EZ-C1 confocal system (Nikon). Two z stacks of the retina for each artery/vein with the capillaries (central and peripheral) were taken and the total number of leucocytes in the arteries, veins and capillaries were counted.

Assessment of retinal glia and microglia Microglial cells were identified using rat anti-mouse F4/80 antigen (1:50; AbD Serotec, Kidlington, UK) on retinal flat mounts. The state of microglial activation was ascertained using previously published methods [17]. Müller cell activation was assessed by staining glial fibrillary acidic protein (GFAP) (1:500; Rabbit anti-GFAP; Dako) on retinal sections from mice that had been diabetic for 24 weeks.

Analysis of retinal DNA damage DNA damage in the retina was determined by TdT-mediated dUTP-X nick end labelling (TUNEL) assay using the In Situ Cell Death Detection Kit (fluorescein; Roche Applied Science, Penzberg, Germany) according to the manufacturer's instructions [15]. Positive cells were assessed by image analysis in multiple sections. Images were taken at three separate points on the central retina at magnification $\times 40$ and presented as the average nuclei in the ganglion cell layer. The slides were viewed using epifluorescent microscopy and analysed using NIS Elements (Nikon).

Quantification of capillary degeneration Retinal flat mounts were prepared for immunofluorescence evaluation as previously described [15]. Briefly, retinas were stained with biotinylated isolectin GS-IB4 (Sigma) and rabbit anti-mouse collagen IV (Col IV) (1:50; AbD Serotec) overnight and imaged using confocal microscopy (Nikon NIS Elements Basic Research (BR) Version 3.0, TE EZ-C1 confocal system). Four regions were taken at magnification $\times 40$ in the central and peripheral retina. Acellular capillaries $<1.3 \mu\text{m}$ were assessed in this study, as in our previous study [15]. Retinal pericytes in the various mouse groups were assessed using the trypsin digest technique as previously described [18].

Graphical presentation and statistical analysis Statistical analysis and graphical presentation was carried out using Prism 5.0 software (Prism 5, San Diego, CA, USA). One-way ANOVA was conducted to compare overall differences among multiple groups and post hoc comparisons were performed using Tukey's or Bonferroni's test. A p value of <0.05 was deemed statistically significant.

Results

Characterisation of diabetes STZ induced a consistent state of diabetes in experimental mice. Analysis of body weight revealed a 14% and 37% reduction in WT and *Rage*^{-/-} diabetic mice compared with age-matched non-diabetic controls at 24 weeks of diabetes (ESM Fig. 1 a, b). HbA_{1c} showed an approximately twofold increase (at 24 weeks) in diabetic mice compared with non-diabetic mice ($p<0.001$; ESM Fig. 1c). Blood glucose was significantly increased in the diabetic mice of both WT and *Rage*^{-/-} mice at 24 weeks of diabetes (ESM Fig. 1d).

MG and its detoxification in the diabetic retina Compared with non-diabetic WT controls, *Rage* mRNA expression in the retina was increased after 12 weeks of diabetes ($p<0.01$) but this differential was not apparent at 24 weeks (ESM Fig. 2). Since AGEs are key ligands for RAGE in the diabetic retina [9] and MG-derived adducts are the most abundant in this tissue during diabetes [19], we examined the accumulation of the reactive AGE precursor MG in the retina and also the activity of its detoxifying enzyme GLO-1. Examination of retinal homogenates showed that after both 8 and 24 weeks of diabetes, WT diabetic mice had significantly higher levels of MG in the retina compared with non-diabetic controls; this elevation was not apparent in *Rage*^{-/-} diabetic mice (Fig. 1 a, b).

Retinal GLO-1 enzymatic activity and mRNA expression for different durations of diabetes was examined. At 4 weeks, retinal GLO-1 activity was significantly enhanced in *Rage*^{-/-} mice, irrespective of the presence of diabetes (Fig. 2a). Not only was GLO-1 activity increased, but also immunofluorescent staining suggested that the expression level of GLO-1 was increased at 8 weeks in *Rage*^{-/-} mouse retinas (Fig. 2b). The transcript of *Glo-1* was also increased significantly at 12 weeks in *Rage*^{-/-} mouse retinas (Fig. 2c).

Since reduced GSH is required as a co-factor for GLO-1 during detoxification of MG [20], we examined the level of reduced GSH in the retinas. Immunofluorescent staining revealed that reduced GSH occurred at higher levels in the retina of *Rage*^{-/-} mice when compared with WT counterparts, irrespective of the presence of diabetes (ESM Fig. 3).

***Rage* deletion protects against retinal vasopermeability** Vasopermeability was assessed 4 weeks after the induction

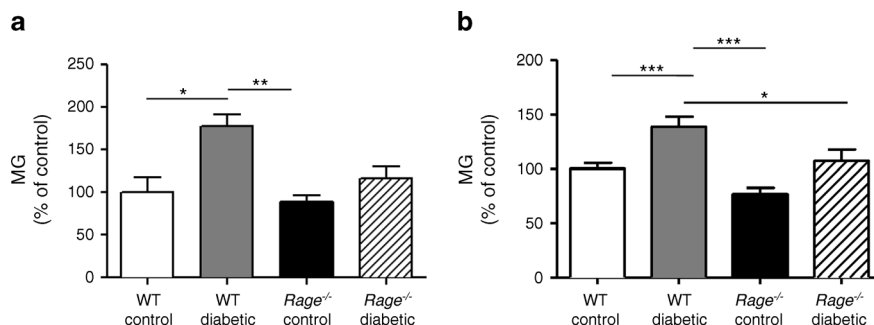


Fig. 1 Accumulation of MG-derived adducts in diabetic mouse retina. MG levels were increased in the retina after both 8 (a) and 24 (b) weeks of diabetes in WT mice. Levels were determined as nmol/mg and presented

as a percentage of control. Data are expressed as mean \pm SEM ($n=3$ per group). * $p<0.05$, ** $p<0.01$ and *** $p<0.001$ for indicated comparisons

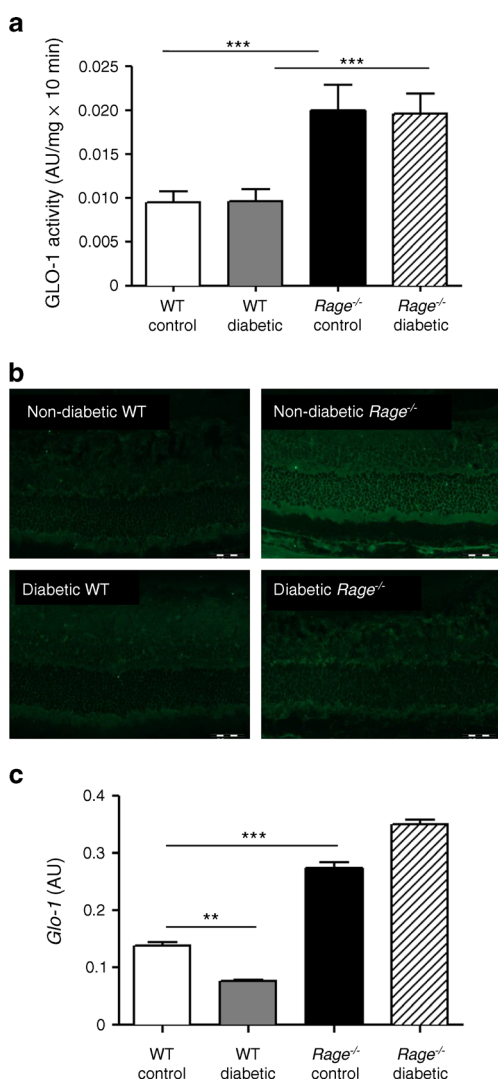


Fig. 2 GLO-1 enzymatic activity in diabetic mouse retina. (a) GLO-1 enzymatic activity is shown in arbitrary units (AU)/mg \times 10 min. (b) GLO-1 immunoreactivity was elevated in non-diabetic *Rage*^{-/-} and diabetic *Rage*^{-/-} mouse retina compared with the low expression in WT control and WT diabetic mouse retina. (c) Retinal *Glo-1* mRNA expression (normalised vs β -actin) in WT diabetic mouse retina was lower than in the WT control and expression in *Rage*^{-/-} control mouse retina was higher. Data are expressed as mean \pm SEM ($n=5$ per group). ** $p<0.01$ and *** $p<0.001$ for the indicated comparisons

of diabetes. This time point was chosen since a significant breakdown of the inner BRB in parallel with capillary leucostasis has been shown in this acute phase [21, 22]. Vasopermeability was assessed through visualisation of albumin that had extravasated and was localised in the neural retina. Albumin remained localised to the vessel lumens of the retinal circulation and beneath the retinal pigment epithelium in non-diabetic WT mice (Fig. 3a). In contrast to their non-diabetic counterparts, diabetic WT mice showed significantly greater areas of focal leakage ($p<0.01$) (Fig. 3a, b). Diabetic *Rage*^{-/-} mice appeared to be protected against albumin leakage into the neural retina since there was no significant difference in area of leakage compared with non-diabetic *Rage*^{-/-} mice (Fig. 3b).

Regulatory role of RAGE in retinal vascular leucostasis Leucostasis is one of the earliest features in murine diabetic retina and can be detected from as early as 1 week after diabetes induction, persisting for at least 3 months [23]. We studied leucostasis after 4 weeks of diabetes using Concanavalin A-FITC to visualise ‘sticking’ leucocytes inside blood vessels. There was a significant increase in the number of adherent leucocytes in the WT diabetic retina (a response that was apparent for arteries, veins and capillaries) when compared with WT non-diabetic controls ($p<0.05$) (Fig. 4a). Diabetic *Rage*^{-/-} mice showed a significant reduction in adherent leucocytes compared with diabetic WT mice (Fig. 4a, b).

***Rage* deletion attenuates diabetes-related glial and neuronal dysfunction** Glial responses were evaluated using the stress-indicator GFAP. In the non-diabetic mouse retina, GFAP was limited in the astrocytes and end feet of Müller cells (Fig. 5a). In diabetic mice GFAP could be detected in the cell processes of Müller cells, indicating Müller cell activation or damage. *Rage*^{-/-} diabetic mice did not show this typical upregulation of GFAP in the Müller cells (Fig. 5a).

TUNEL-positive cells, indicating DNA strand breaks, were significantly more abundant in the diabetic mice than in non-diabetic controls ($p<0.001$) (Fig. 5b). TUNEL-positive cells were mainly localised to the ganglion cell layer but they were also present in the inner and outer nuclear layers. In *Rage*^{-/-}

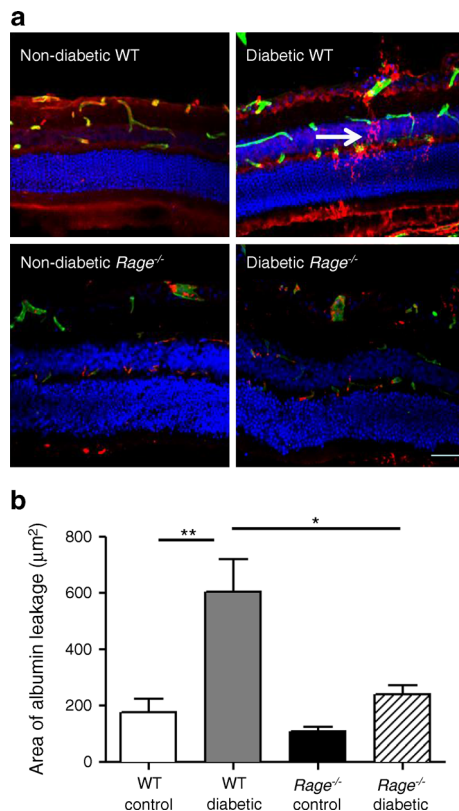


Fig. 3 Vasopermeability in diabetic mouse retina is modulated by *Rage*. (a) Blood vessel leakage was assessed by albumin (red) staining after 4 weeks of diabetes. There was a significant increase in albumin that had leaked from the blood vessels (stained green with isolectin B4) in WT diabetic mice compared with the control WT mice (white arrow showing leakage). (b) Graph of albumin leakage (area in μm^2) from the retinal blood vessels of *Rage*^{-/-} diabetic mice showed a decrease in the amount of leakage compared with that in WT diabetic mice. Data are expressed as mean \pm SEM ($n=6$). * $p<0.05$ and ** $p<0.01$ for the indicated comparisons

mice, diabetic conditions also increased the number of TUNEL-positive cells, especially in the ganglion cell layer ($p<0.01$) (Fig. 5b). However, comparison between the WT and *Rage*^{-/-} diabetic mice showed that the absence of RAGE significantly protected the diabetic retina against DNA breaks ($p<0.05$) (Fig. 5b).

RAGE plays a role in microglial activation during diabetic retinopathy An increase in microglial activation and number of cells in the neuropile has been previously demonstrated in diabetic retinopathy [24]. Following both 12 and 24 weeks diabetes duration in WT mice, there was a significant increase in the numbers of F4/80-positive microglia in the retina when compared with non-diabetic controls ($p<0.001$) (Fig. 6a). *Rage*^{-/-} mice did not demonstrate this diabetes-induced increase and, indeed, there were significantly less positive microglia in the retinas of *Rage*^{-/-} diabetic mice compared with WT diabetic mice ($p<0.001$) (Fig. 6a).

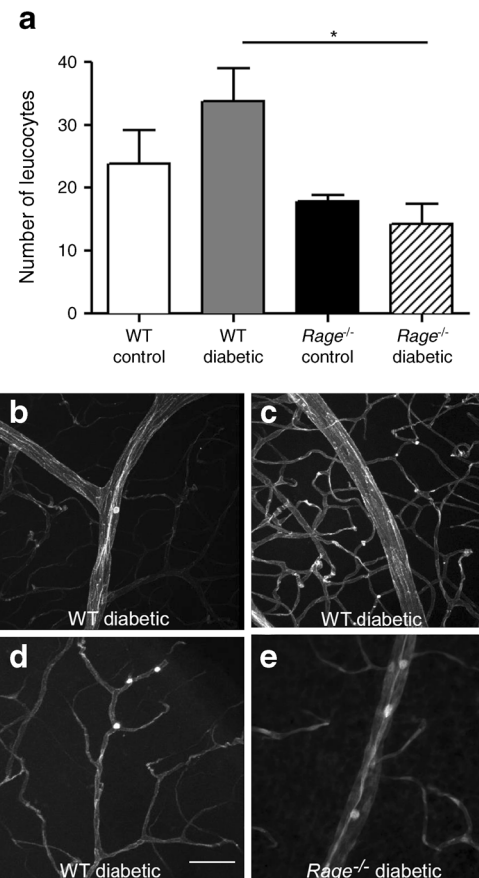


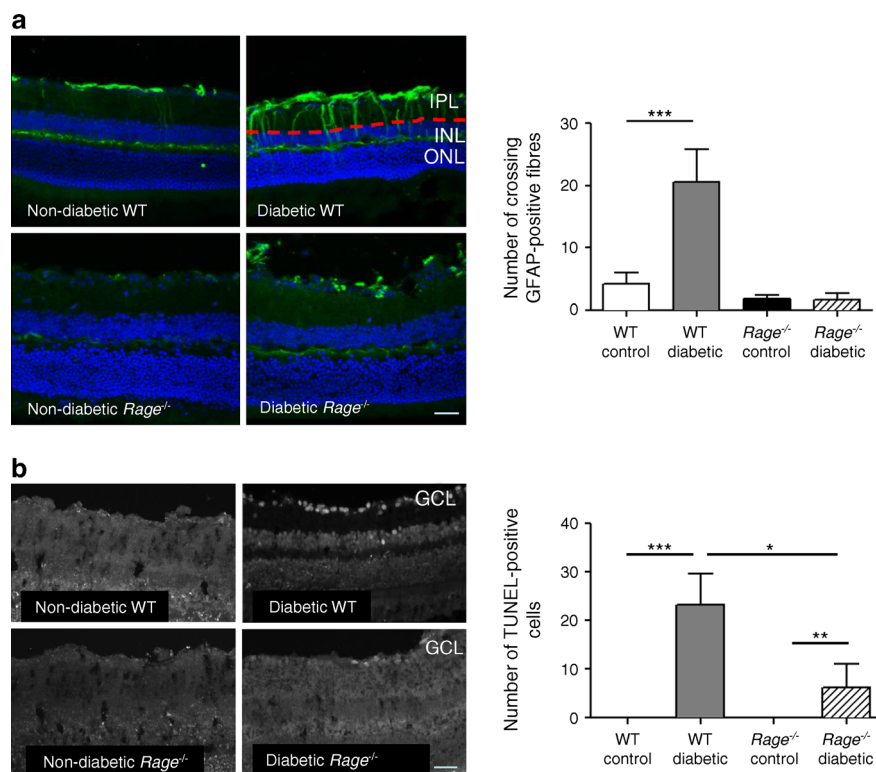
Fig. 4 Role of *Rage* deletion in retinal vascular leucostasis during diabetes. (a) Leucostasis was assessed in the mouse retina after 4 weeks of diabetes covering 0.1 mm^2 as the mean of 16 z stacks per retina. Data are expressed as mean \pm SEM ($n=6$). * $p<0.05$ for indicated comparison. Leucocytes are shown: at the branches in the vein of a WT diabetic mouse retina (b); in the capillaries of a WT diabetic mouse retina (c); at the junctions of the capillaries of a WT diabetic mouse retina (d) and in the vein of a *Rage*^{-/-} diabetic mouse retina (e). Scale bar, $50\text{ }\mu\text{m}$

Microglia presented in various states of activation in normal retina from both WT and *Rage*^{-/-} mice; however, the ‘classical’ resting (ramified/dendritic; Fig. 6b) state was predominant (Fig. 6b, c). Following 12 weeks of diabetes, there was a significant increase in activated microglia in the WT diabetic retina whereas this increase was absent in *Rage*^{-/-} mice (Fig. 6c). After 24 weeks of diabetes, there was no apparent increase in microglial activation compared with non-diabetic controls, only an increase in resting microglia between WT control and WT diabetic (Fig. 6d).

RAGE deletion protects against diabetes-mediated retinal capillary degeneration Acellular capillary formation and the death of pericytes and endothelium in the retina is a hallmark in mice beyond 5 months of diabetes [25]. By definition, acellular capillaries do not have endothelial cells and the vessels are isolectin B4 negative but since a basement membrane persists in these acellular vessels, Col IV immunoreactivity remains present (Fig. 7a, b). When retinal flat mounts were

Fig. 5 *Rage* deletion alters GFAP expression in retinal Müller glia and DNA strand breaks in ganglion cells. (a)

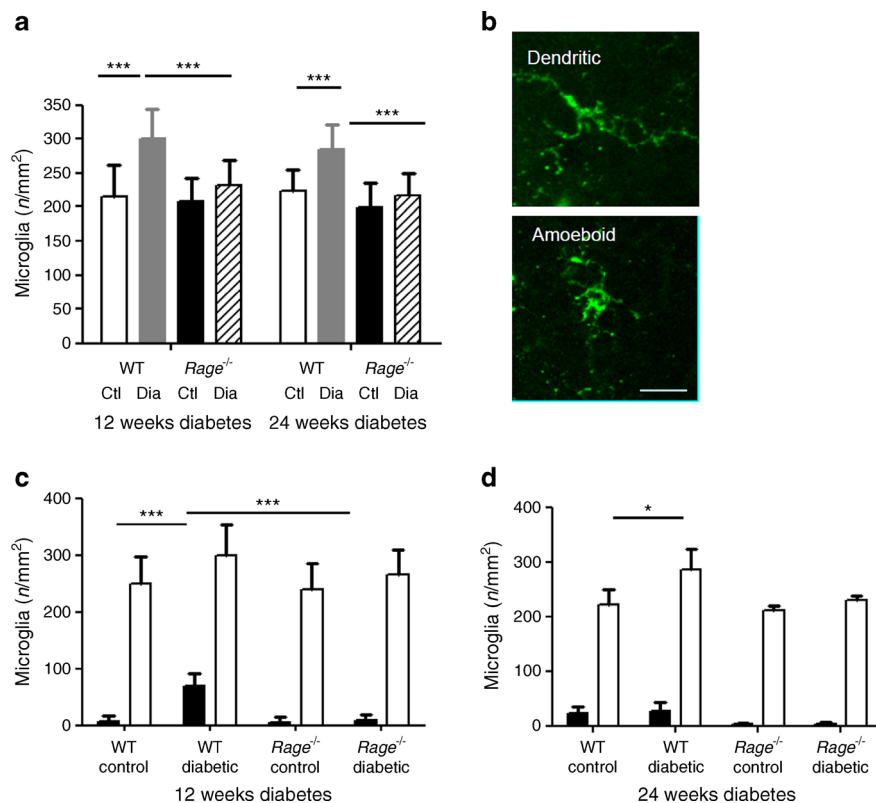
Immunostaining of GFAP and graph showing the mean GFAP-positive fibres crossing the inner plexiform layer (IPL) and the inner nuclear layer (INL) in the cohort. The red dotted line indicates the border between the IPL and the INL where the GFAP-positive fibres cross. ONL, outer nuclear layer. (b) TUNEL positivity was assessed on the sections of retina from all the experimental groups. GCL, ganglion cell layer. Scale bar, 50 μ m. Data are expressed as mean \pm SEM ($n=4$). * $p<0.05$, ** $p<0.01$ and *** $p<0.001$ for indicated comparisons



evaluated and quantified for isolectin/Col IV staining, there was a significant increase in acellular capillaries in WT diabetic mice compared with non-diabetic controls ($p<0.001$)

(Fig. 7c). This diabetes-mediated pathology was absent in the retina from *Rage*^{-/-} diabetic mice. In trypsin-digested preparations, the number of pericytes in the retinal capillaries

Fig. 6 Microgliosis in mouse retinas at 12 and 24 weeks of diabetes. (a) Total microglia cell numbers at 12 and 24 weeks of diabetes illustrating more microglia in the WT diabetic mouse than in the other groups. Ctl, control; Dia, diabetic. (b) Representative images of dendritic (resting) and amoeboid (active) microglia, taken from *Rage*^{-/-} control groups. Scale bar, 25 μ m. (c, d) Activation states of microglia were ascertained, post image capture. These graphs illustrate activated (black bars) and resting microglia (white bars) labelled with F4/80 after 12 and 24 weeks of diabetes. Data are expressed as mean \pm SEM ($n=6$). * $p<0.05$ and *** $p<0.001$ for indicated comparisons



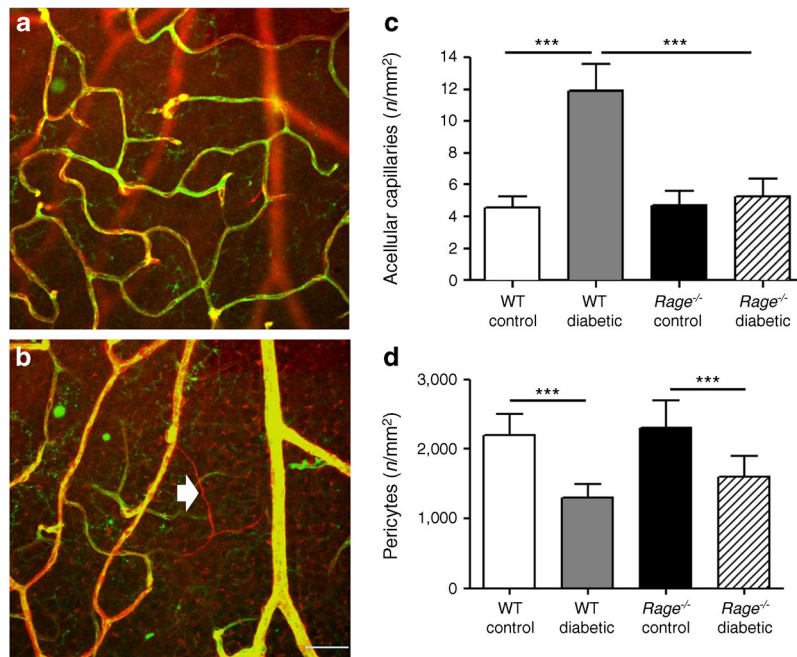


Fig. 7 Retinal capillary degeneration and pericyte dropout during diabetes after 24 weeks of diabetes. Flat-mounted retina from WT control and diabetic mice in the deeper plexus (**a**) and the superficial plexus (**b**), illustrating capillaries visualised using isolectin B4 (endothelium; green) and Col IV (basement membrane; red). Acellular capillaries show continuance of Col IV positivity but loss of endothelium (arrow). Scale bar,

50 μ m. (**c**) Graph showing the mean number of acellular capillaries in the retina. Data are expressed as mean \pm SEM ($n=6$). *** $p<0.001$ for the indicated comparisons. (**d**) Graph showing pericyte coverage in the retina. Data expressed as mean \pm SD ($n=7$). *** $p<0.001$ for the indicated comparisons

was significantly reduced in diabetic WT mice compared with controls. Pericyte loss was also significant in the *Rage*^{-/-} diabetic mice (Fig. 7d).

Discussion

RAGE plays a significant role in inflammation-linked disorders such as Alzheimer's disease and atherosclerosis [26, 27]. In the context of diabetic complications, activation of RAGE by a range of ligands has been strongly associated with inflammatory pathology in the kidney and peripheral nerves and vessels [28]. In the current investigation, we show that *Rage*^{-/-} mice are protected from some of the key early-stage lesions of diabetic retinopathy.

Rage deletion had no significant impact on glycaemia (based on HbA_{1c}) or characteristic deficits in weight gain. However, diabetes did increase RAGE expression in the murine retina after 12 weeks of diabetes, although this elevation was not maintained up to 24 weeks. RAGE expression is widespread in the retinal neuropile of rats and mice, being especially high in the Müller glia [4], and it has been shown to increase in diabetic rats concomitantly with ligands such as AGEs and S100b [6, 29].

AGEs are ligands for RAGE and many are derived from the α -oxoaldehyde MG, which occurs at high

levels in the diabetic retina and gives rise to adducts such as N^ε-(carboxy-ethyl) lysine and MG-derived hydroimidazolone-1 [30]. An important defence against MG-related toxicity is the glyoxalase complex (formed from GLO-1 and glyoxalase II components), which converts MG to D-lactate using GSH as a co-factor [31]. We have previously reported that overexpression of GLO-1 in diabetic rats protects against retinopathy [30]. The current study now shows that *Rage*^{-/-} mice possess significantly enhanced retinal GLO-1 activity, a higher GLO-1 expression level and enhanced GSH levels when compared with WT mice. This explains the reduced levels of MG in *Rage*^{-/-} diabetic mice compared with their WT diabetic counterparts. The relationship between RAGE and GLO-1 during diabetes has been reported previously [32, 33] and it has been suggested that upregulation of the enzyme is linked to RAGE-mediated transcriptional activation of Myd88 [34]. In the current study we have demonstrated that *Rage*^{-/-} mice have higher levels of retinal GSH than their WT counterparts and this may also account for why GLO-1 activity is also elevated in these mice.

Interactions between circulating immune cells and the retinal vasculature probably begin with leucostasis, which may contribute to capillary occlusion and breakdown of the BRB [35]. An earlier study suggested that neutrophils become less sticky after treatment with RAGE antibody [36]. It is possible that leucocytes from *Rage*^{-/-} diabetic mice are less sticky due to the absence of this receptor. Deletion of *Rage* in diabetic

mice provided protection against vascular leucostasis and vasopermeability and this agrees with the findings of previous studies that used therapeutic approaches. For example, reducing the bioavailability of RAGE ligands (using sRAGE) prevented vasopermeability in mice [12]. Deletion of *Rage* transcription resulted in a marked protection against adhesion of leucocytes. While it is uncertain whether the acute-phase leucostasis phenomenon is a major contributor to capillary degeneration [37], it seems likely that pro-inflammatory processes involving bone marrow-derived cells make a contribution to diabetic retinopathy [38]. Although not evaluated in the current study, there is evidence that TLRs may be involved in retinopathy [39]. The established interplay between TLRs and RAGE is potentially important and could contribute to innate immune responses in the retina, especially since they share common ligand interactions with S100 and HMGB1 [10]. Müller glia act as inflammatory activators and this response is diminished by sRAGE treatment in *db/db* mice [4] and, as shown in the current study, genetic deletion of *Rage*. Professional immune cells, which reside in proximity to retinal blood vessels (perivascular macrophages) or within various layers of the neuropile (microglia), are of even more critical importance [40]. While these cells have a homeostatic function, they are also linked to neuroinflammation in the human diabetic retina [24] and animal models [41, 42]. RAGE plays a role in monocyte migration and microglial activation [43, 44] by altering the expression of adhesion molecules, facilitating monocytic extravasation into the CNS and altering microglial cells from resting to activated states [45].

The current study suggests a role for RAGE in microglial activation, especially after 12 weeks of diabetes at which time *Rage*^{−/−} mice were protected from this response. This bimodal response according to the duration of diabetes may also reflect the altered expression levels of RAGE between 12 and 24 weeks. Further supporting evidence comes from *Rage*^{−/−} mice subjected to laser-induced choroidal neovascularisation; these mice showed reduced lesion size alongside diminution of immune cell activation when compared with WT controls [46].

In rodent models of diabetic retinopathy, pericyte dropout and acellular capillary formation are ‘gold standard’ markers of retinopathy progression. In the current study, analysis using both trypsin digests and confocal microscopy of whole mounts showed significant vasodegeneration at 24 weeks in WT diabetic mice. *Rage*^{−/−} mice were protected from diabetes-induced acellular capillary formation, providing strong evidence of an important role for this receptor in diabetic retinopathy. The fact that pericyte loss was not markedly prevented when RAGE was absent is in keeping with studies using RAGE blockade [38] and suggests that loss of these cells is not strongly associated with the RAGE pathway. Our data support the suggestion that inflammatory cascades are linked to retinal vascular damage as diabetes progresses. As

a component of the innate immune response, RAGE plays an important role in diabetic retinopathy.

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Contribution statement CMV was involved with the conception and design of experiments, researched data and wrote the manuscript. MW, LMC, JGF TF, AS and MK researched data and contributed to the manuscript. AB provided *Rage*^{−/−} mice. HPH was involved with the conception and design of some experiments and contributed to the manuscript. MC researched and interpreted data and wrote some of the manuscript. AWS conceived and designed experiments, interpreted data, wrote the manuscript and guided the project. All authors revised the paper for important intellectual content and approved the final version.

AWS is the guarantor of this work and, as such, has full access to all data in the study and takes responsibility for the integrity of data and accuracy of the data analysis.

Duality of interest The authors declare that there is no duality of interest associated with this manuscript.

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